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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/573,332	04/06/2007	David W. Morris	PP023362.0003	5041
27476 NOVARTIS V	7590 03/03/201 ACCINES AND DIAC		EXAM	IINER
INTELLECTU	JAL PROPERTY- X10	HARRIS, ALANA M		
P.O. BOX 809 Emeryville, C.			ART UNIT	PAPER NUMBER
• /-			1643	
			MAIL DATE	DELIVERY MODE
			03/03/2010	PAPER

Please find below and/or attached an Office communication concerning this application or proceeding.

The time period for reply, if any, is set in the attached communication.

# Application No. Applicant(s) MORRIS ET AL. 10/573,332 Office Action Summary Examiner Art Unit Alana M. Harris, Ph.D. 1643 -- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --Period for Reply A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) OR THIRTY (30) DAYS. WHICHEVER IS LONGER, FROM THE MAILING DATE OF THIS COMMUNICATION. Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication. If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication - Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b). Status 1) Responsive to communication(s) filed on 17 November 2009. 2a) ☐ This action is FINAL. 2b) This action is non-final. 3) Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under Ex parte Quayle, 1935 C.D. 11, 453 O.G. 213. Disposition of Claims 4) Claim(s) 1-55 is/are pending in the application. 4a) Of the above claim(s) 1-32 and 37-55 is/are withdrawn from consideration. 5) Claim(s) \_\_\_\_\_ is/are allowed. 6) Claim(s) 33-36 is/are rejected. 7) Claim(s) \_\_\_\_\_ is/are objected to. 8) Claim(s) \_\_\_\_\_ are subject to restriction and/or election requirement. Application Papers 9) The specification is objected to by the Examiner. 10) The drawing(s) filed on is/are; a) accepted or b) objected to by the Examiner. Applicant may not request that any objection to the drawing(s) be held in abevance. See 37 CFR 1.85(a). Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d). 11) The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152. Priority under 35 U.S.C. § 119 12) Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f). a) All b) Some \* c) None of: Certified copies of the priority documents have been received. 2. Certified copies of the priority documents have been received in Application No. Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)). \* See the attached detailed Office action for a list of the certified copies not received.

1) Notice of References Cited (PTO-892)

Paper No(s)/Mail Date 04/10/2008.

2) Notice of Draftsperson's Patent Drawing Review (PTO-948)
3) Information Disclosure Statement(s) (PTO/S5/08)

Attachment(s)

Interview Summary (PTO-413)
 Paper No(s)/Mail Date.

6) Other:

5) Notice of Informal Patent Application

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#### DETAILED ACTION

## Election/Restrictions

1. Applicant's election with traverse of Group VIII (claims 33 and 34) in the reply filed on November 17, 2010 is acknowledged. The traversal is on the ground(s) that "Groups VIII and IX (claims 35 and 36) are both directed to embodiments that comprise at least two polynucleotides" and consequently a search of both sets of claims would be duplicative and "...should not be an undue burden on the Examiner". This is found persuasive and both Groups will be examined.

The remainder of the requirement is still deemed proper and is therefore made FINAL.

2. Claims 1-55 are pending.

Claims 1-32 and 37-55, drawn to non-elected inventions are not examined on the merits.

Claims 33-36 are examined on the merits.

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# Claim Rejections - 35 USC § 112

3. The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

4. Claims 35 and 36 are rejected under 35 U.S.C. 112, second paragraph, as

being indefinite for failing to particularly point out and distinctly claim the

subject matter which applicant regards as the invention.

a. Claims 35 and 36 are drawn to an electronic library. It is not clear

if Applicants intended to state an electronic medium, an array or microarray

that comprises sequences. The term, electronic library has not been defined in

the specification. Accordingly, the metes and bounds cannot be determined.

### Claim Rejections - 35 USC § 102

The following is a quotation of the appropriate paragraphs of 35
 U.S.C. 102 that form the basis for the rejections under this section made in this
 Office action:

A person shall be entitled to a patent unless -

(e) the invention was described in (1) an application for patent, published under section 122(b), by another filed in the United States before the invention by the applicant for patent or (2) a patent granted on an application for patent by another filed in the United States before the invention by the applicant for patent, except that an international application filed under the treaty defined in section 351 (a) shall have the effects for purposes of this subsection of an application filed in the United States only if the international application designated the United States and was published under Article 21(2) of such treaty in the English language.

(e) the invention was described in a patent granted on an application for patent by another filed in the United States before the invention thereof by the applicant for patent, or on an international application by another who has fulfilled the requirements of paragraphs (1), (2), and (4) of section 371(c) of this title before the invention thereof by the applicant for patent.

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The changes made to 35 U.S.C. 102(e) by the American Inventors

Protection Act of 1999 (AIPA) and the Intellectual Property and High Technology

Technical Amendments Act of 2002 do not apply when the reference is a U.S.

patent resulting directly or indirectly from an international application filed

before November 29, 2000. Therefore, the prior art date of the reference is

determined under 35 U.S.C. 102(e) prior to the amendment by the AIPA (pre
AIPA 35 U.S.C. 102(e)).

6. Claims 33 and 35 are rejected under 35 U.S.C. 102(e) as being anticipated by Knoll et al./ U.S. Patent number 7,014,997 B2 (filed May 14, 2001). Knoll discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; column 1, lines 36-45; and column 14, lines 56-65. Knoll discloses sequences 425 and 148, which are the same as Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, respectively, corresponding hybridization probes, arrays, and kits in which these components are contained, see abstract; column 2, lines 5-50; column 3, lines 17-25; bridging paragraph of columns 24 and 25; and the claims beginning in column 37. These components are useful in detecting "[c]hromosomal abnormalities often common and ...diagnostic in...leukemia and other cancers". see column 2. 1st full sentence.

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In absence of a clear definition of the term, electronic library as cited in the pending 112, 2<sup>nd</sup> paragraph rejection, the art reads on claim 35. The Examiner regards the cited arrays as the equivalent of Applicants' electronic library.

Search results between Applicants' SEQ ID NO: 4 and Knoll sequence 425.

```
US-09-854-867-425, rni
; Sequence 425, Application US/09854867
: Patient No. 7014997
: GENERAL INFORMATION:
; APPLICANT: JOAN, KNOLL H
; APPLICANT: ROGAN, PETER K
; TITLE OF INVENTION: SINGLE COPY GENOMIC HYBRIDIZATION PROBES AND METHOD OF GENERATING SAME
; FILE REFERENCE: 30307
  CURRENT APPLICATION NUMBER: US/09/854,867
; CURRENT FILING DATE: 2003-05-08
; NUMBER OF SEQ ID NOS: 613
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 425
  LENGTH: 3285
   TYPE: DNA
   ORGANISM: Homo sapiens
  FEATURE:
 Query Match
                        24.1%; Score 289.6; DB 5; Length 3285;
 Best Local Similarity 65.9%;
 Matches 567; Conservative 5; Mismatches 216; Indels 72; Gaps
          34 CAAAGAATAAGAGAAACACATAAAGCAAGAAATAAGAGAAACACATAATTCAACAATAA 93
Db
         365 CAAAATACATAAAGCAAATACTAATAGAACTGAAAAGAGAAATAGACAAATCCACAATAA 424
          94 TA----GAAACTTAAATATCCTCCTTTCAATAATAGATACAACAACTAAGCAGTTGATCA 149
         425 TAGTTGGAGACTTCAATACCCCACTGTCAGTAATTGATAGATCAACTAGACAGAAAATCA 484
         150 ACAAGAAAACAGAAGATTTGAACAATGCTATGAACCAACTAGACCTAACGTCTA---TCT 206
         485 ATAAGGATATAGAAGACTTGAACAACACTATCAACCAACTGGACCTAATTGACATATTAT 544
QУ
         207 ATAAAACACCACCCAACAACAACAGCAGAATACATATTCTTCTCAGATATACATAGAACATTC 266
         545 AGAACACTCCACCCAACAACAGCAGAATACACATTCTTYTCAAGTGCACATGGAACATTC 604
         267 TCCAGGATAGGCCATCTGTTAGGACATAAAACAAGTCTCAAAAAATGTAAAAGAATTGAG 326
Db
         605 ACCAAGATAGACCATATGCTRGGCCATAAAACAAGTCTCAATAAATTTAAAAAAATTGAA 664
         327 ATCAGACAAAGTCTGTTCTCTGACCACAA------CCAGTAACAGAAG 368
         665 ATYATACAAAGTATGTTCTCTGACCACAATGGAATAAAAWTAGAAATCAATAACAAAAG 724
         369 GAAATTTGAAGAATCCATAAGTATGTGGAAATGAATCAAGGAACT----- 413
         725 ATACTCTGGAAAATNCACAAATACTTGGAAATTAAACAACATACTTCTAAATAACTCATG 784
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Qу	414	CAAGGGAAATTAGAAAATACTTTGAAATGAATGAAAATGAAAA	456
Db	785	$\tt GGTCAAAGAAATCAAAAGAGAAATTAAAAAATATTTTGAAATAAAT$	844
Qу	457	CACAGCATACCAAAAC-TTATGAGATGCAGCTAAAATAGTGCTTACAGAGAAATTAATAG	515
Dlo	845	YACAACATATCAAAAATTTATGGGATGCAGCTAAAGCAGTGCTTAGAGGGAAATTTATAG	904
Qу	516	$\tt CTATTAATGCCTGTATTTTTAAAAGAAGAAGAAGATACCAAATTAAAAAAAA$	575
dd	905	CATTAAATGCCTATATTAAAAAGAAGAAGATCTCAAATCAATAACCTAAGTTTCCA	962
Qу	576	TTTAAGAAAAAGAATAGTGAACCAAGCCCAAATCAAGCAGAAGGAAG	627
Db	963	CCTTAAGAAACTAGAAAAAGAAGAGCAAATTAAACCCAAAGTAAGCAGAAGAAAGGAAAG	1022
Qy	628	${\tt AATAAAGATTAGAATGGAAAAAATGAAATATGGAATTGGAAAAACTAGAGAAAAATTAA}$	687
Db	1023	AATAAAGATTAGAGCAGAAATAAATGAAATAGAAAACAGAAAAACAATAGAAAAATCAA	1082
Qу	688	$\tt CAAACCCAAAAGTTGTTATATCAAAAAGATTGATAAGTTTGATAAACATTTAACTAGACT$	747
Db	1083	TAAAACCAAAAGTTGGTTCTTTGAAAAGATAAAATTGACAAACCTTTAGCTAGACT	1138
Qу	748	TACCCTAATATCAAAACCACATACAGATATCACAAGAAAAGTACAGACCAATATCTCTCA	807
Db	1139	AAAAAAAGGGGGAGACACAAATTACTAATANATCAGAAATGAAAGGGGGGAYATTACT	1198
Qу	808	TAAGACACATATAAGATAGA 827	
Db	1199	ACAGATYCTACAGAAATAAA 1218	

# Search results between Applicants' SEQ ID NO: 427 and Knoll sequence 148.

	ult No.	Score	Query Match	Length	DB	ID	Description
	1	1200	100.0	76360	16	US-10-669-920-427	Sequence 427, App
	2	1200	100.0	76360	19	US-10-573-332-427	Sequence 427, App
	3	939	78.2	75853	7	US-10-087-192-382	Sequence 382, App
C	4	506.6		665590	18	US-10-990-328-94275	Sequence 94275, A
C	5	484.6		161994	24	US-11-112-908-57	Sequence 57, Appl
C	6	484.6		303956	27	US-11-033-056A-36950	Sequence 36950, A
C	7	483	40.2	29142	10	US-10-741-600-17977	Sequence 17977, A
C	8	483	40.2		17	US-10-796-280-12569	Sequence 12569, A
C	9	483	40.2		36	US-12-287-505-17977	Sequence 17977, A
C	10	483	40.2		36	US-12-337-905-17977	Sequence 17977, A
	11	473.8	39.5	130877	9	US-10-322-281-54	Sequence 54, Appl
	1.2	473.8	39.5	130877	16	US-10-539-228-54	Sequence 54, Appl
C	13	464.8	38.7	102139	18	US-10-990-328-95533	Sequence 95533, A
	1.4	464.8	38.7	439596	27	US-11-033-056A-37873	Sequence 37873, A
C	15	460.4	38.4	50498	17	US-10-105-299-9098	Sequence 9098, Ap
C	16	460.4	38.4	50498	18	US-10-868-184-6947	Sequence 6947, Ap
C	17	448.2	37.4		18	US-10-990-328-96764	Sequence 96764, A
	18	446.2		330955	36	US-12-113-481-90	Sequence 90, Appl
	19	445	37.1	340000	24	US-11-102-978-3	Sequence 3, Appli
C	20	444.6	37.1	103000	27	US-11-033-056A-38751	Sequence 38751, A
C	21	441	36.8	2418	3	US-09-854-867-148	Sequence 148, App
c	22	441	36.8	2418	11	US-10-786-970A-148	Sequence 148, App

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7. Claims 33 and 35 are rejected under 35 U.S.C. 102(e) as being anticipated by Cargill et al./ U.S. Patent Application Publication number 2005/0026169 A1 (effective filing date April 30, 2003). Cargill discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; page 3, sections 0023 and 0024; and page 14, section 0153. Cargill discloses sequences 17996 and 17977, which are the same as Applicants' SEQ ID NO: 4 and SEQ ID NO: 427, respectively, corresponding hybridization probes, arrays, DNA chips and kits in which these components are contained, see abstract; page 14, sections 0154 and 0155; page 15, section 0162; and page 16, section 0173-page 18, section 0186.

In absence of a clear definition of the term, electronic library as cited in the pending 112, 2<sup>nd</sup> paragraph rejection the art reads on claim 35. The Examiner regards the cited arrays and chips as the equivalent of Applicants' electronic library.

## Search results between Applicants' SEQ ID NO: 427 and Cargill sequence 17977.

```
US-10-741-600-17977, rnpbm
; Sequence 17977, Application US/10741600
: Publication No. US20050026169A1
; GENERAL INFORMATION:
; APPLICANT: CARGILL, Michele et al.
; TITLE OF INVENTION: GENETIC POLYMORPHISMS ASSOCIATED WITH
; TITLE OF INVENTION: MYOCARDIAL INFARCTION, METHODS OF DETECTION AND USES THEREOF; FILE REFERENCE; CL001499
: CURRENT APPLICATION NUMBER: US/10/741,600
; CURRENT FILING DATE: 2003-12-22
; NUMBER OF SEQ ID NOS: 73997
; SOFTWARE: FastSEQ for Windows Version 4.0
: SEO ID NO 17977
   LENGTH: 29142
   TYPE: DNA
  ORGANISM: Homo sapiens
US-10-741-600-17977
 Query Match
                      40.2%; Score 483; DB 10; Length 29142;
 Best Local Similarity 73.0%;
 Matches 788; Conservative 0; Mismatches 215; Indels 76; Gaps 10;
         130 AAAAATACAAAAATTAGCTGGGTGTGGTGGTGCATGTCTGTAATCCCAGCTACTCAGGAG 189
       26298 AAAAATATTTTTTTTGGCCGAGTGCGGTGGCTCATGCCTGTAATCCCAGCTACTCAGGAG 26239
         190 GCTGAGGCTTGAACCTGG-----GAGTCAGAGGTTGCAATGAGCCGAGATCGCAC 239
       26238 GCTGAGGCAGGAGAATCGCTTGAACCTAGAAGCAGAGGTTGCAGTGAGCTGAGATCATGC 26179
         240 CACTGCACTCCAGCCTGGC------GACAGAGCAAGACTCCTTGTCAAAA-AAAAAA 290
       26178 CACGGCACTCCAGCCTGGCCGACAGAGCCGACGAAGGAGTTGGTCTCAACAGTGGCTTAA 26119
Οv
         291 AAAATTCAGTAAACCATACTGTAAACAGATGTGCTGTCATTCAGGCTTAGTTATGCCATT 350
       26118 AATGTTCGGTAAACCATGCTGGGAACAGATGGGCTGTCATCCAGGCTTTGTTGTTCCATC 26059
         351 TACTGAACACAAGCAGAGTAGATTTAGCATAATTTCTAACAGCAATAGGATTTTTGAAAT 410
QУ
       26058 CACAGAGCACAGGCAGAATAGATTTAGCATCATTCTTAAGAGCCCTGGGATTTTCGGAAT 25999
         411 GGTAAATGAACACTGGCTTCGACTTAAACTCACCAGCTGCATTACCTCCTAACAAGAGAG 470
       25998 GGTAAATGAGCATTGGCTTCAACTTAAAGTCGCCAGCTGCATTAGC----CCTAACGAG 25944
         471 AGTCAGCCTGTCCTTTGAAGCTCTGAAGCCAGACATTGACTT---ATCTGTAGCTAGAAA 527
       25943 AGTCAGCCTGTCCTTCGAAGCTTTGAAGCCAGGGATTGACTTCTCTTCTGTAGCTATGAA 25884
         25883 AGTCCCAGGTGGCATCTTCTTCCAATAGAAGGCTGTTTGTCTACATTGAAAGTCTGTTTT 25824
         579 -----AAATCTATGATCTTAGCTAGATCTTCTGGATGACTTGCTGCAGCT 623
       25823 TGAGTGTAGCCACCTTCGTCAGTGAGATCAGCTAGATCTGATGGATAACTTACTGCAGCT 25764
         624 TCTACATCAGCACTTGCTGCTTCACCTTGCACCTTTATTATATGGAGATGGCTTGTTTCC 683
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Db
       25703 TTCAACCTCACAAAACAACCTCTGTTAGCTTCTGACTTTTCTTCTGCAACTTTCTCACCT 25644
         740 CCCTCAGCCTTCACAGAATTGAAGAGTTAGGGCATTGCTATGGATTAGGCTTTGGCTT 799
        25643 CTCTCTGCCTTCAAAGGATTGAAGAGTGAGGGCCTTTCTCCGGGACTAGGCTTTGGCTT 25584
         800 AAGGGAATGTTATGGCTGGTTTGATCTTCTATCCAGATCACCCAAACTTTCCTCCATATC 859
       25583 AAAGGAATGTTGTGGCTGATTTGATCTTCTTTCCAGA----CTCAAACTTTCTCCGTATC 25528
         860 AGCAATAATGCTGTTTTGCTTTTTTGTCATTCATATGTTCACTGGAGTAGCACTTTTAAT 919
        25527 AGCAATAATGCTATTCTGCTTTCATATCATTTTGTGTGTCCACTAGAGCAGCACTTTTAAT 25468
         920 T-----TTTTTTTTTTTCCTTGATACAGAGTCTCGCTCTGTCACCCAGGC 963
        25467 TTAATTTATCATTTTTATTTTTTTTTTGGGACAGAATTTCGCTCTGTCACCCAGGT 25408
         964 TGGAGTGCAATGGCGTGGTCTCGGCTCACTGCAACCTCAGCCTCCTGGGTTCAAGTGGTT 1023
       25407 TGGAGTGCAGTGGTGCGATCTTGACTCATGGTAACCTCCGCCTCCCCACTTCAAGCGATT 25348
        1024 CTCCCGCCTCAGCCTCCAGAGTAGCTGGGGCTACAGGGGGCGCCACCACCACACCCAGC-CA 1082
        25347 CTCCTGCCTCAGCGTCCTAAGTAGTTCGGAGTACAGGCATGCACCACCACGCCCAGCTAA 25288
        1083 GTTTTTGTATTTTTAGTAGACACGGGGTTTCACTATGTTGGCCAGGCTGGACTCGAACT 1141
       25287 TTTTTTGTATTTTAGTAGAGATGGGGTCTTGCCATGTTGGCCAGGCTGGTCTCGAACT 25229
```

## rni. Search results between Applicants' SEQ ID NO: 4 and Cargill sequence 17996.

Result		Query					
No.	Score	Match	Length	DB	ID	Description	
1	289.6	24.1	3285	3	US-09-573-080A-425	Sequence 425, App	
2	289.6	24.1	3285	5	US-09-854-867-425	Sequence 425, App	
3	245.8	20.5	278866	3	US-09-949-016-13922	Sequence 13922, A	
4	245.8	20.5	278866	3	US-09-949-016-13923	Sequence 13923, A	
5	245.8	20.5	278866	3	US-09-949-016-13924	Sequence 13924, A	
6	245.8	20.5	278866	3	US-09-949-016-13925	Sequence 13925, A	
7	245.8	20.5	278866	3	US-09-949-016-13926	Sequence 13926, A	
8	245.8	20.5	278866	3	US-09-949-016-14699	Sequence 14699, A	
9	245.8	20.5	278866	3	US-09-949-016-14700	Sequence 14700, A	
10	245.8	20.5	278866	3	US-09-949-016-14701	Sequence 14701, A	
1.1	245.8	20.5	278866	3	US-09-949-016-14702	Sequence 14702, A	
1.2	245.8	20.5	278866	3	US-09-949-016-14703	Sequence 14703, A	
13	232.8	19.4	999	10	US-10-301-480C-341210	Sequence 341210,	
1.4	220.6	18.4	148783	3	US-09-949-016-15729	Sequence 15729, A	
1.5	215.2	17.9	998	10	US-10-301-480C-367650	Sequence 367650,	
1.6	215.2	17.9	999	10	US-10-301-480C-267544	Sequence 267544,	
1.7	215.2	17.9	999	10	US-10-301-480C-367649	Sequence 367649,	
18	212.2	17.7	7678	3	US-09-573-080A-348	Sequence 348, App	
19	212.2	17.7	7678	5	US-09-854-867-348	Sequence 348, App	
c 20	211	.8	17.7	416	65 8 US-10-741-600	0-17996 Sequence 17996,	A

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8. Claims 34 and 36 are rejected under 35 U.S.C. 102(e) as being anticipated by Venter et al./ U.S. Patent number 6,812,339 B1 (filed September 10, 2001). Venter discloses sequences that would selectively hybridize to Applicants' SEQ ID NO: 53 and SEQ ID NO: 622, fragments thereof or their complements within in a sample, see sequence alignment information following instant rejection; column 5, lines 32-47; and column 18, lines 24-63. Venter discloses sequences 340 and 3171, which are the same as Applicants' SEQ ID NO: 53 and SEQ ID NO: 622, respectively, corresponding hybridization probes, arrays, DNA chips, computer-based and data storage systems and kits in which these components are contained, see column 14, lines 39-54; and columns 15 and 16.

In absence of a clear definition of the term, electronic library as cited in the pending 112,  $2^{nd}$  paragraph rejection, the art reads on claim 35. The Examiner regards the cited arrays, chips and systems as the equivalent of Applicants' electronic library.

### Search results between Applicants' SEQ ID NO: 53 and Venter sequence 340.

```
RESULT 3
; Sequence 340, Application US/09949016
: Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
  TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
: FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
: CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
: PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
  PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 340
  LENGTH: 1145
   TYPE: DNA
   ORGANISM: Human
US-09-949-016-340
 Query Match 50.1%; Score 305.4; DB 3; Length 1145; Best Local Similarity 99.7%; Pred. No. 8.2e-75;
 Matches 306; Conservative 0; Mismatches 1; Indels 0; Gaps
         95 AGCAGGACAGGCTGCTTTGGTTTGTGACCTCCAGGCAGGACGCCATCCTCTCCAGAATG 154
         79 AGCAGGACAGGCTGCTTTGGTTTGTGACCTCCAGGCAGGACGGCCATCCTCTCCAGAATG 138
Db
        215 ATGTGCTTCTCCTGCTTGAACCAGAAGAGCAATCTGTACTGCCTGAAGCCGACCATCTGC 274
        199 ATGTGCTTCTCCTGCTTGAACCAGAAGAGCAATCTGTACTGCCTGAAGCCGACCATCTGC 258
        275 TCCGACCAGGACAACTACTGCGTGACTGTGTCTGCTAGTGCCGGCATTGGGAATCTCGTG 334
        259 TCCGACCAGGACAACTACTGCGTGACTGTGTCTGCTAGTGCCGGCATTGGGAATCTCGTG 318
        335 ACATTTGGCCACAGCCTGAGCAAGACCTGTTCCCCGGCCTGCCCCATCCCAGAAGGCGTC 394
        319 ACATTTGGCCACAGCCTGAGCAAGACCTGTTCCCCGGCCTGCCCATCCCAGAAGGCGTC 378
Qν
        395 AATGTGG 401
Db
        379 AATGTTG 385
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Search results between Applicants' SEQ ID NO: 622 and Venter sequences 3171 and 3172.

	ult No.	Score		Length		ID	Description
	1	3270.8				US-10-648-593-132	Sequence 132, App
	2	3270.8	79.0	4574	8	US-11-072-175-132	Sequence 132, App
	3	2841.2	68.6	4268	3	US-09-954-556-3	Sequence 3, Appli
	4	2525.8	61.0	3416	2	US-08-451-822A-15	Sequence 15, Appl
	5	2525.8	61.0	3416	3	US-08-323-430-15	Sequence 15, Appl
C	6	2425.6	58.6	4367	5	US-10-021-698A-3641	Sequence 3641, Ap
	7	2413.8		3025	3	US-09-954-556-23	Sequence 23, Appl
	8	2335		3080	3	US-09-954-556-25	Sequence 25, Appl
	9	2269		3106		US-10-701-263-1	Sequence 1, Appli
		2199.2		3244		US-09-954-556-24	Sequence 24, Appl
G	11	2199.2		3244	5	US-10-021-698A-3626	Sequence 3626, Ap
		1957		2310		US-08-471-570-9	Sequence 9, Appli
		1949.6		2923		US-09-954-556-20	Sequence 20, Appl
		1944.4		2826		US-09-954-556-21	Sequence 21, Appl
		1943		2868		US-09-954-556-19	Sequence 19, Appl
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		1834.4		2650		US-09-954-556-28	Sequence 28, Appl
		1834.4		2676		US-08-471-570-7	Sequence 7, Appli
		1688.4				US-09-954-556-10	Sequence 10, Appl
	20	1583			2	US-08-471-570-3	Sequence 3, Appli
		1460.4		1954		US-08-471-570-5	Sequence 5, Appli
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	24	1332.8	32.2				Sequence 3173, Ap

9. Any inquiry concerning this communication or earlier communications from the Examiner should be directed to Alana M. Harris, Ph.D. whose telephone number is (571)272-0831. The Examiner works a flexible schedule, however she can normally be reached Monday through Saturday, 7:30 am to 6:30 pm with alternate Fridays off.

If attempts to reach the Examiner by telephone are unsuccessful, the Examiner's supervisor, Larry R. Helms, Ph.D. can be reached on (571) 272-0832. The fax phone number for the organization where this application or proceeding is assigned is 571-273-8300.

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Alana M. Harris, Ph.D. 26 February 2010 /Alana M. Harris, Ph.D./ Primary Examiner, Art Unit 1643